

ACTGTCTGGAAGCTGGACTGAGTCACCAAAAGGCGAATGGCTTCATCTTATAAAATGTCTGAACAAAGCACA  
ACTTCTGAGCACATTTTACAGAAGACATGTGATCACCTGATCCTGACTAACCGTTCTGGATTAGAGACAGA  
CTCAGTAGCAGAGGAAATGAAGCAGACTGTGGAGGGACAGGGGCATACAGTGCAGTGGGCAGCTCTCCTGA  
TACTCGCGGTGATAATACCCACCATTGGTGGGAACATCCTTGTGATTCTGGCTGTTGCACTGGAGAAAAGG  
CTGCAGTACGCTACCAACTACTTTTTTAATGTCTTGGCGATAGCAGATTTGCTGGTTGGATTGTTTGTGAT  
GCCGATTGCCCTCTTGACAATCATGTTTGAGGCTATATGGCCCCTCCCAGTGGCCCTGTGTCCTGCCTGGT  
TATTCCTCGATGTTCTCTTTTCAACTGCCTCCATCATGCATCTCTGTGCCATTTCCCTGGACCGCTATATA  
GCCATCAAAAAGCCAATTCAGGCCAATCAGTGCAACACCCGGGCTACTGCATTCATCAAGATTACAGTGGT  
ATGGTTAATTTCAATAGGCATCGCCATCCCAGTCCCTATTAAAGGAATCGAGACTGATGTGATTAATCCAC  
ACAATGTCACCTGTGAGCTGACAAAGGACCGCTTTGGCAGTTTTATGGTCTTTGGGTCACTGGCTGCTTTC  
TTCGTACCTCTCACCATCATGGTAGTCACTTACTTTCTCACCATTACACTTTACAGAAGAAAGCTTACTT  
GGTCAAAAATAAGCCACCTCAACGCCTAACACGGTGGACTGTGCCCACAGTTTTCTTAAGGGAAGACTCAT  
CCTTTTCATCACCAGAAAAGGTGGCAATGCTGGATGGGTCTCACAGGGATAAAATTCTACCTAACTCAAGT  
GATGAGACACTTATGCGAAGAATGTCCTCAGTTGGAAAAAGATCAGCCCAAACCATTTCTAATGAGCAGAG  
AGCCTCGAAGGCCCTTGGAGTCGTGTTTTCTCTTTTCTGCTTATGTGGTGCCCTTTTTTTATTACAAATC  
TAACTTTAGCTCTGTGTGATTCTTGCAATCAGACCACTCTCAAAACACTCCTGGAGATATTTGTGTGGATA  
GGCTACGTTTTCTCGGGGGTGAATCCTCTGATCTATACACTCTTCAATAAGACATTTTCGGGAAGCATTTGG  
CAGGTACATCACCTGCAATTACCGAGCCACAAAGTCAGTAAAAGCACTTAGGAAGTTTTCCAGTACACTTT  
GTTTTGGGAATTCAATGGTAGAAAACCTCTAAATTTTTTCACAAAACATGGAATTCGAAATGGGATCAACCT  
GCCATGTACCAGAGCCCAATGAGGCTCCGATGTTCAACCATTAGTCCTCATCAATCATCCTCCTCGATAC  
CCTTCTCACTGAAAACGATGGCGACAAAGCGGAAGAGCAGGTCAGCTACATATTGCAGGAACGGGCGGCC  
TCATCTTGAGAGAGGGTGATGAGCAGGACGCGCACCATGGCAGGTTCAAGAGTGA (SEQ ID NO:1)

MASSYKMSEQSTTSEHILQKTC DHLILTNRS GLETD SVAEEMKQTVEGQGHTVHWAALLILAVI IPTIGGN  
ILVILAVALEKRLQYATNYFLMSLA IADLLVGLFVMPIALLTIMFEAIWPLPLALCPAWLFLDVLFSTASI  
MHLCAISLDRYIAIKKPIQANQCNTRATAFIKITV VWLISIGIAIPVPIKGIETDVINPHNVTCELT KD RF  
GSFMVFGSLAAFFVPLTIMVVTYFLTIHTLQKKAYLVKNKPPQRLTRWTVPTVFLREDSSFSSPEKVAMLD  
GSHRDKILPNSSDETLMRMSSVGKRS AQTISNEQRASKALGVVFFLFLLMWCPFFITNLTLALCDSCNQ T  
TLKTLLEIFVWIGYVSSGVNPLIYTLFNKTFREAFGRYITCN YRATKSVKALRKFSSTLCFGNSMVENS KF  
FTKHGIRNGINPAMYQSPMRLRCSTIQSSSIILLDTLLTENDGDKAE EQVS YILQERAGLILREGDEQDAR  
APWQVQE (SEQ ID NO:2)

FIGURE 1

Underlined = deleted in targeting construct

**Bold** = sequence flanking Neo insert in targeting construct

**ACTGTCTGGA****ACTGG****ACTGAGT****CACCAAAAGGCGAATGGCTTCATCTTATAAAATGTCTG**  
**AACAAAGCACAACTTCTGAGCACATTTTACAGAAGACATGTGATCACCTGATCCTGACTA**  
**ACCGTTCTG**GATTAGAGACAGACTCAGTAGCAGAGGAAATGAAGCAGACTGTGGAGGGAC  
AGGGGCATACAGTGCACTGGGCAGCTCTCCTGATACTCGCGGTGATAATACCCACCATTG  
GTGGGAACATCCTTGTGATTCTGGCTGTTGCACTGGAGAAAAGGCTGCAGTACGCTACCA  
ACTACTTTTAAATGTCCTT **GCGGATAGCAGATTTGCTGGTTGGATTGTTTGTGATGCCGA**  
**TTGCCCTCTTGACAATCATGTTTG****AGGCTATATGGCCCCCTCCCACTGGCCCTGTGTCCTG**  
**CCTGGTTATTCTCGATGTTCTCTTTTCAACTGCCTCCATCATGCATCTCTGTGCCATTT**  
**CCCTGGACCGCTATATAGCCATCAAAAAGCCAATTCAGGCCAATCAGTGCAACACCCGGG**  
**CTACTGCATTTCATCAAGATTACAGTGGTATGGTTAATTTCAATAGGCATCGCCATCCCAG**  
**TCCCTATTAAAGGAATCGAGACTGATGTGATTAATCCACACAATGTCACCTGTGAGCTGA**  
**CAAAGGACCGCTTTGGCAGTTTATGGTCTTTGGGTCACCTGGCTGCTTTCTTCGTACCTC**  
**TCACCATCATGGTAGTCACTTACTTTCTCACCATTACACTTTTACAGAAGAAAGCTTACT**  
**TGGTCAAAAATAAGCCACCTCAACGCCTAACACGGTGGACTGTGCCCACAGTTTTCTCTAA**  
**GGGAAGACTCATCCTTTTCATCACCAGAAAAGGTGGCAATGCTGGATGGGTCTCACAGGG**  
**ATAAAATTCTACCTAACTCAAGTGATGAGACACTTATGCGAAGAATGTCCTCAGTTGGAA**  
**AAAGATCAGCCCCAAACCATTTCTAATGAGCAGAGAGCCTCGAAGGCCCTTGGAGTCGTGT**  
**TTTTCTTTTTCTGCTTATGTGGTGCCCCCTTTTTTATTACAAATCTAACTTTAGCTCTGT**  
**GTGATTCCTGCAATCAGACCACTCTCAAAACACTCCTGGAGATATTTGTGTGGATAGGCT**  
**ACGTTTCCTCGGGGGTGAATCCTCTGATCTATACACTCTTCAATAAGACATTTTCGGGAAG**  
**CATTTGGCAGGTACATCACCTGCAATTACCGAGCCACAAAGTCAGTAAAAGCACTTAGGA**  
**AGTTTTCCAGTACACTTTGTTTTGGGAATTCAATGGTAGAAAACCTCTAAATTTTTACAA**  
**AACATGGAATTCGAAATGGGATCAACCCTGCCATGTACCAGAGCCCAATGAGGCTCCGAT**  
**GTTCAACCATTCAGTCCTCATCAATCATCCTCCTCGATACCTTCTCACTGAAAACGATG**  
**GCGACAAAGCGGAAGAGCAGGTCAGCTACATATTGCAGGAACGGGCCGCTCATCTTGA**  
**GAGAGGGTGATGAGCAGGACGCACGCGCACCATTGGCAGGTTCAAGAGTGA**

FIGURE 2A

Gene Sequence  
Structure \*

130 bp

Sequence Deleted

319 bp

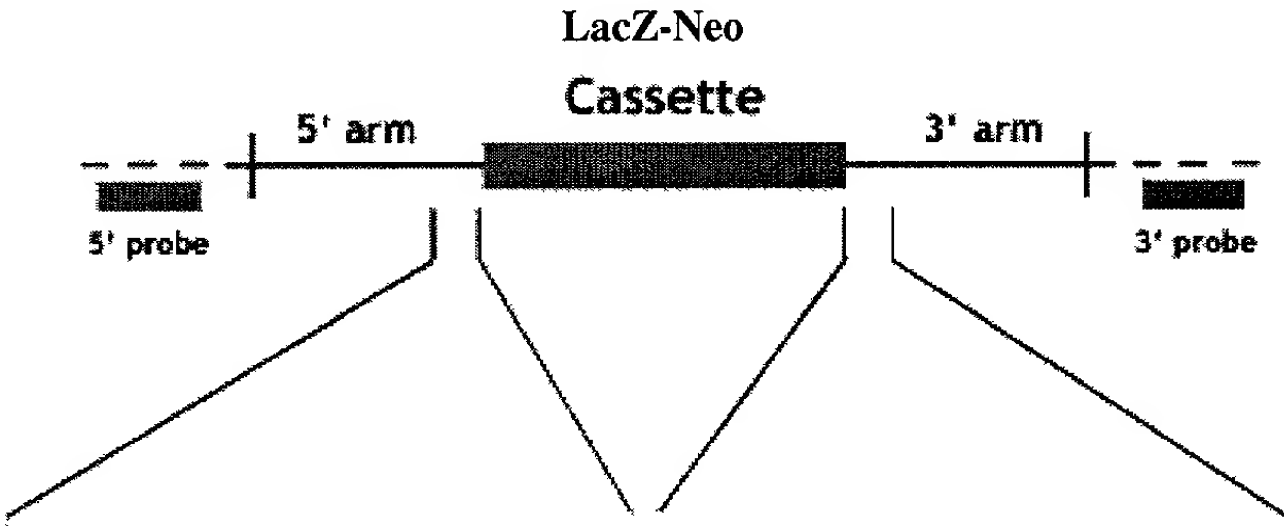
Size of full-length  
cDNA: 1550 bp



Targeting Vector\* (genomic sequence)

Construct Number: 2520

Arm Length:  
5': 1.6 kb  
3': 5 kb



————— Targeting Vector  
- - - - - Endogenous Locus

\* Not drawn to scale

<b>5'</b> >TGAGTGTCTGGTGGGTTTGCT AAATGCTTTGCTAAAGCAGATGAC TTGCTTAGCTACTGACCATGCTGA CCACTGTCTGGAAGTGGACTGAGT CACCAAAAGGCGAATGGCTTCATC TTATAAAATGTCTGAACAAAGCAC AACTTCTGAGCACATTTTACAGAA GACATGTGATCACCTGATCCTGAC TAACCGTTCTG<3' (SEQ ID NO:3)	<b>5'</b> >GGCGATAGCAGATTGCTGGT TGGATTGTTTGTGATGCCGATTGC CCTCTTGACAATCATGTTTGGTGA GTATTTCCCCTTGTTCCCTGCCACT GAACACTACTAACGTAGTGAAATG GACACTCACTGACCTTTATTTTGT TTGAAATAAAAGAAGGACCTGGAT TAAAAACACAGAAGGGAACATTCC TTCATTTTTC<3' (SEQ ID NO:4)
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FIGURE 2B